

=====

Sequence Listing was accepted.

See attached Validation Report.

If you need help call the Patent Electronic Business Center at (866)
217-9197 (toll free).

Reviewer: Keisha Douglas

Timestamp: [year=2008; month=6; day=3; hr=19; min=19; sec=55; ms=600;]

=====

Application No: 10550778 Version No: 1.0

Input Set:

Output Set:

Started: 2008-05-05 17:01:41.676
Finished: 2008-05-05 17:02:05.718
Elapsed: 0 hr(s) 0 min(s) 24 sec(s) 42 ms
Total Warnings: 25
Total Errors: 25
No. of SeqIDs Defined: 25
Actual SeqID Count: 25

Error code	Error Description
W 213	Artificial or Unknown found in <213> in SEQ ID (1)
E 257	Invalid sequence data feature in <221> in SEQ ID (1)
W 213	Artificial or Unknown found in <213> in SEQ ID (2)
E 257	Invalid sequence data feature in <221> in SEQ ID (2)
W 213	Artificial or Unknown found in <213> in SEQ ID (3)
E 257	Invalid sequence data feature in <221> in SEQ ID (3)
W 213	Artificial or Unknown found in <213> in SEQ ID (4)
E 257	Invalid sequence data feature in <221> in SEQ ID (4)
W 213	Artificial or Unknown found in <213> in SEQ ID (5)
E 257	Invalid sequence data feature in <221> in SEQ ID (5)
W 213	Artificial or Unknown found in <213> in SEQ ID (6)
E 257	Invalid sequence data feature in <221> in SEQ ID (6)
W 213	Artificial or Unknown found in <213> in SEQ ID (7)
E 257	Invalid sequence data feature in <221> in SEQ ID (7)
W 213	Artificial or Unknown found in <213> in SEQ ID (8)
E 257	Invalid sequence data feature in <221> in SEQ ID (8)
E 257	Invalid sequence data feature in <221> in SEQ ID (8)
W 213	Artificial or Unknown found in <213> in SEQ ID (9)
E 257	Invalid sequence data feature in <221> in SEQ ID (9)
W 213	Artificial or Unknown found in <213> in SEQ ID (10)

Input Set:

Output Set:

Started: 2008-05-05 17:01:41.676
Finished: 2008-05-05 17:02:05.718
Elapsed: 0 hr(s) 0 min(s) 24 sec(s) 42 ms
Total Warnings: 25
Total Errors: 25
No. of SeqIDs Defined: 25
Actual SeqID Count: 25

Error code	Error Description
E 257	Invalid sequence data feature in <221> in SEQ ID (10)
W 213	Artificial or Unknown found in <213> in SEQ ID (11)
E 257	Invalid sequence data feature in <221> in SEQ ID (11)
W 213	Artificial or Unknown found in <213> in SEQ ID (12)
E 257	Invalid sequence data feature in <221> in SEQ ID (12)
W 213	Artificial or Unknown found in <213> in SEQ ID (13)
E 257	Invalid sequence data feature in <221> in SEQ ID (13)
W 213	Artificial or Unknown found in <213> in SEQ ID (14)
E 257	Invalid sequence data feature in <221> in SEQ ID (14)
W 213	Artificial or Unknown found in <213> in SEQ ID (15)
E 257	Invalid sequence data feature in <221> in SEQ ID (15)
W 213	Artificial or Unknown found in <213> in SEQ ID (16)
E 257	Invalid sequence data feature in <221> in SEQ ID (16)
W 213	Artificial or Unknown found in <213> in SEQ ID (17)
E 257	Invalid sequence data feature in <221> in SEQ ID (17)
W 213	Artificial or Unknown found in <213> in SEQ ID (18)
E 257	Invalid sequence data feature in <221> in SEQ ID (18)
W 213	Artificial or Unknown found in <213> in SEQ ID (19)
E 257	Invalid sequence data feature in <221> in SEQ ID (19) This error has occurred more than 20 times, will not be displayed
W 213	Artificial or Unknown found in <213> in SEQ ID (20) This error has occurred more than 20 times, will not be displayed

<110> Zumbrunn, Jurg
 Lociuero, Sergio
 Gombert, Frank
 Moehle, Kerstin
 DeMarco, Steven J.
 Vrijbloed, Jan Wim
 Mukherjee, Reshmi
 Obrecht, Daniel
 Romagnoli, Barbara
 Robinson, John Anthony

<120> Template-Fixed Peptidomimetics as Medicaments against HIV and
 Cancer

<130> 753-54 PCT/US

<140> 10550778
 <141> 2008-05-05

<150> PCT/EP2003/04641
 <151> 2003-05-02

<160> 25

<170> PatentIn version 3.4

<210> 1
 <211> 12
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Template-fixed peptidomimetic

<220>
 <221> MISC_FEATURE
 <222> (3)..(3)
 <223> Xaa = 2-Nal

<220>
 <221> DISULFID
 <222> (4)..(11)

<220>
 <221> MISC_FEATURE
 <222> (7)..(7)
 <223> Xaa = D-Lys

<220>
 <221> MISC_FEATURE
 <222> (10)..(10)
 <223> Xaa = Cit

<220>

<221> MOD_RES

<222> (12)..(12)

<223> AMIDATION

<400> 1

Arg Arg Xaa Cys Tyr Arg Xaa Pro Tyr Xaa Cys Arg

1 5 10

<210> 2

<211> 12

<212> PRT

<213> Artificial Sequence

<220>

<223> Template-fixed peptidomimetic

<220>

<221> MISC_FEATURE

<222> (3)..(3)

<223> Xaa = 2-Nal

<220>

<221> DISULFID

<222> (4)..(11)

<220>

<221> MISC_FEATURE

<222> (7)..(7)

<223> Xaa = D-Lys

<220>

<221> MISC_FEATURE

<222> (10)..(10)

<223> Xaa = Cit

<220>

<221> MOD_RES

<222> (12)..(12)

<223> AMIDATION

<400> 2

Arg Arg Xaa Cys Tyr Lys Xaa Pro Tyr Xaa Cys Arg

1 5 10

<210> 3

<211> 12

<212> PRT

<213> Artificial Sequence

<220>

<223> Template-fixed peptidomimetic

<220>
<221> MISC_FEATURE
<222> (3)..(3)
<223> Xaa = 2-Nal

<220>
<221> DISULFID
<222> (4)..(11)

<220>
<221> MISC_FEATURE
<222> (7)..(7)
<223> Xaa = D-Lys

<220>
<221> MISC_FEATURE
<222> (10)..(10)
<223> Xaa = Cit

<220>
<221> MOD_RES
<222> (12)..(12)
<223> AMIDATION

<400> 3

Arg Arg Xaa Cys Tyr Lys Xaa Pro Arg Xaa Cys Arg
1 5 10

<210> 4
<211> 12
<212> PRT
<213> Artificial Sequence

<220>
<223> Template-fixed peptidomimetic

<220>
<221> MISC_FEATURE
<222> (3)..(3)
<223> Xaa = 2-Nal

<220>
<221> DISULFID
<222> (4)..(11)

<220>
<221> MISC_FEATURE
<222> (7)..(7)
<223> Xaa = D-Pro

<220>
<221> MOD_RES

<222> (12)..(12)

<223> AMIDATION

<400> 4

Arg Arg Xaa Cys Tyr Lys Xaa Pro Tyr Arg Cys Arg
1 5 10

<210> 5

<211> 12

<212> PRT

<213> Artificial Sequence

<220>

<223> Template-fixed peptidomimetic

<220>

<221> MISC_FEATURE

<222> (3)..(3)

<223> Xaa = 2-Nal

<220>

<221> DISULFID

<222> (4)..(11)

<220>

<221> MISC_FEATURE

<222> (7)..(7)

<223> Xaa = D-Pro

<220>

<221> MOD_RES

<222> (12)..(12)

<223> AMIDATION

<400> 5

Arg Arg Xaa Cys Tyr Arg Xaa Pro Tyr Arg Cys Arg
1 5 10

<210> 6

<211> 12

<212> PRT

<213> Artificial Sequence

<220>

<223> Template-fixed peptidomimetic

<220>

<221> MISC_FEATURE

<222> (3)..(3)

<223> Xaa = 2-Nal

<220>
<221> DISULFID
<222> (4)..(11)

<220>
<221> MISC_FEATURE
<222> (10)..(10)
<223> Xaa = Cit

<220>
<221> MOD_RES
<222> (12)..(12)
<223> AMIDATION

<400> 6

Arg Arg Xaa Cys Tyr Arg Lys Pro Tyr Xaa Cys Arg
1 5 10

<210> 7
<211> 14
<212> PRT
<213> Artificial Sequence

<220>
<223> Template-fixed peptidomimetic

<220>
<221> MISC_FEATURE
<222> (3)..(3)
<223> Xaa = 2-Nal

<220>
<221> DISULFID
<222> (4)..(13)

<220>
<221> MISC_FEATURE
<222> (6)..(6)
<223> Xaa = Cit

<220>
<221> MISC_FEATURE
<222> (8)..(8)
<223> Xaa = D-Pro

<220>
<221> MISC_FEATURE
<222> (12)..(12)
<223> Xaa = Cit

<220>
<221> MOD_RES
<222> (14)..(14)
<223> AMIDATION

<400> 7

Arg Arg Xaa Cys Tyr Xaa Lys Xaa Pro Tyr Arg Xaa Cys Arg
1 5 10

<210> 8

<211> 14

<212> PRT

<213> Artificial Sequence

<220>

<223> Template-fixed peptidomimetic

<220>

<221> MOD_RES

<222> (1)..(1)

<223> ACETYLATION

<220>

<221> MISC_FEATURE

<222> (3)..(3)

<223> Xaa = 2-Nal

<220>

<221> DISULFID

<222> (4)..(13)

<220>

<221> MISC_FEATURE

<222> (6)..(6)

<223> Xaa = Cit

<220>

<221> MISC_FEATURE

<222> (8)..(8)

<223> Xaa = D-Pro

<220>

<221> MISC_FEATURE

<222> (12)..(12)

<223> Xaa = Cit

<220>

<221> MOD_RES

<222> (14)..(14)

<223> AMIDATION

<400> 8

Arg Arg Xaa Cys Tyr Xaa Lys Xaa Pro Tyr Arg Xaa Cys Arg
1 5 10

<210> 9

<211> 14
<212> PRT
<213> Artificial Sequence

<220>
<223> Template-fixed peptidomimetic

<220>
<221> MISC_FEATURE
<222> (1)..(1)
<223> Xaa = iPrArg

<220>
<221> MISC_FEATURE
<222> (3)..(3)
<223> Xaa = 2-Nal

<220>
<221> DISULFID
<222> (4)..(13)

<220>
<221> MISC_FEATURE
<222> (6)..(6)
<223> Xaa = Cit

<220>
<221> MISC_FEATURE
<222> (8)..(8)
<223> Xaa = D-Pro

<220>
<221> MISC_FEATURE
<222> (12)..(12)
<223> Xaa = Cit

<220>
<221> MOD_RES
<222> (14)..(14)
<223> AMIDATION

<400> 9

Xaa Arg Xaa Cys Tyr Xaa Lys Xaa Pro Tyr Arg Xaa Cys Arg
1 5 10

<210> 10
<211> 14
<212> PRT
<213> Artificial Sequence

<220>
<223> Template-fixed peptidomimetic

<220>
<221> MISC_FEATURE
<222> (1)..(1)
<223> Xaa = D-Arg

<220>
<221> MISC_FEATURE
<222> (3)..(3)
<223> Xaa = 2-Nal

<220>
<221> DISULFID
<222> (4)..(13)

<220>
<221> MISC_FEATURE
<222> (6)..(6)
<223> Xaa = Cit

<220>
<221> MISC_FEATURE
<222> (8)..(8)
<223> Xaa = D-Pro

<220>
<221> MISC_FEATURE
<222> (12)..(12)
<223> Xaa = Cit

<220>
<221> MOD_RES
<222> (14)..(14)
<223> AMIDATION

<400> 10

Xaa Arg Xaa Cys Tyr Xaa Lys Xaa Pro Tyr Arg Xaa Cys Arg
1 5 10

<210> 11
<211> 14
<212> PRT
<213> Artificial Sequence

<220>
<223> Template-fixed peptidomimetic

<220>
<221> DISULFID
<222> (4)..(13)

<220>
<221> MISC_FEATURE
<222> (6)..(6)
<223> Xaa = Cit

<220>
<221> MISC_FEATURE
<222> (8)..(8)
<223> Xaa = D-Pro

<220>
<221> MISC_FEATURE
<222> (12)..(12)
<223> Xaa = Cit

<220>
<221> MOD_RES
<222> (14)..(14)
<223> AMIDATION

<400> 11

Arg Arg Trp Cys Tyr Xaa Lys Xaa Pro Tyr Arg Xaa Cys Arg
1 5 10

<210> 12
<211> 14
<212> PRT
<213> Artificial Sequence

<220>
<223> Template-fixed peptidomimetic

<220>
<221> MISC_FEATURE
<222> (3)..(3)
<223> Xaa = F(pNH2)

<220>
<221> DISULFID
<222> (4)..(13)

<220>
<221> MISC_FEATURE
<222> (6)..(6)
<223> Xaa = Cit

<220>
<221> MISC_FEATURE
<222> (8)..(8)
<223> Xaa = D-Pro

<220>
<221> MISC_FEATURE
<222> (12)..(12)
<223> Xaa = Cit

<220>
<221> MOD_RES

<222> (14)..(14)

<223> AMIDATION

<400> 12

Arg Arg Xaa Cys Tyr Xaa Lys Xaa Pro Tyr Arg Xaa Cys Arg
1 5 10

<210> 13

<211> 14

<212> PRT

<213> Artificial Sequence

<220>

<223> Template-fixed peptidomimetic

<220>

<221> MISC_FEATURE

<222> (3)..(3)

<223> Xaa = W(6-C1)

<220>

<221> DISULFID

<222> (4)..(13)

<220>

<221> MISC_FEATURE

<222> (6)..(6)

<223> Xaa = Cit

<220>

<221> MISC_FEATURE

<222> (8)..(8)

<223> Xaa = D-Pro

<220>

<221> MISC_FEATURE

<222> (12)..(12)

<223> Xaa = Cit

<220>

<221> MOD_RES

<222> (14)..(14)

<223> AMIDATION

<400> 13

Arg Arg Xaa Cys Tyr Xaa Lys Xaa Pro Tyr Arg Xaa Cys Arg
1 5 10

<210> 14

<211> 14

<212> PRT

<213> Artificial Sequence

<220>
<223> Template-fixed peptidomimetic

<220>
<221> MISC_FEATURE
<222> (1)..(1)
<223> Xaa = (EA)G

<220>
<221> MISC_FEATURE
<222> (3)..(3)
<223> Xaa = 2-Nal

<220>
<221> DISULFID
<222> (4)..(13)

<220>
<221> MISC_FEATURE
<222> (6)..(6)
<223> Xaa = Cit

<220>
<221> MISC_FEATURE
<222> (8)..(8)
<223> Xaa = D-Pro

<220>
<221> MISC_FEATURE
<222> (12)..(12)
<223> Xaa = Cit

<220>
<221> MOD_RES
<222> (14)..(14)
<223> AMIDATION

<400> 14

Xaa Arg Xaa Cys Tyr Xaa Lys Xaa Pro Tyr Arg Xaa Cys Arg
1 5 10

<210> 15
<211> 14
<212> PRT
<213> Artificial Sequence

<220>
<223> Template-fixed peptidomimetic

<220>
<221> MISC_FEATURE
<222> (1)..(1)

<223> Xaa = (PrA)G

<220>

<221> MISC_FEATURE

<222> (3)..(3)

<223> Xaa = 2-Nal

<220>

<221> DISULFID

<222> (4)..(13)

<220>

<221> MISC_FEATURE

<222> (6)..(6)

<223> Xaa = Cit

<220>

<221> MISC_FEATURE

<222> (8)..(8)

<223> Xaa = D-Pro

<220>

<221> MISC_FEATURE

<222> (12)..(12)

<223> Xaa = Cit

<220>

<221> MOD_RES

<222> (14)..(14)

<223> AMIDATION

<400> 15

Xaa Arg Xaa Cys Tyr Xaa Lys Xaa Pro Tyr Arg Xaa Cys Arg

1

5

10

<210> 16

<211> 14

<212> PRT

<213> Artificial Sequence

<220>

<223> Template-fixed peptidomimetic

<220>

<221> MISC_FEATURE

<222> (1)..(1)

<223> Xaa = (BA)G

<220>

<221> MISC_FEATURE

<222> (3)..(3)

<223> Xaa = 2-Nal

<220>

<221> DISULFID
<222> (4)..(13)

<220>
<221> MISC_FEATURE
<222> (6)..(6)
<223> Xaa = Cit

<220>
<221> MISC_FEATURE
<222> (8)..(8)
<223> Xaa = D-Pro

<220>
<221> MISC_FEATURE
<222> (12)..(12)
<223> Xaa = Cit

<220>
<221> MOD_RES
<222> (14)..(14)
<223> AMIDATION

<400> 16

Xaa Arg Xaa Cys Tyr Xaa Lys Xaa Pro Tyr Arg Xaa Cys Arg
1 5 10

<210> 17
<211> 14
<212> PRT
<213> Artificial Sequence

<220>
<223> Template-fixed peptidomimetic

<220>
<221> MISC_FEATURE
<222> (1)..(1)
<223> Xaa = (EGU)G

<220>
<221> MISC_FEATURE
<222> (3)..(3)
<223> Xaa = 2-Nal

<220>
<221> DISULFID
<222> (4)..(13)

<220>
<221> MISC_FEATURE
<222> (6)..(6)
<223> Xaa = Cit

<220>
<221> MISC_FEATURE
<222> (8)..(8)
<223> Xaa = D-Pro

<220>
<221> MISC_FEATURE
<222> (12)..(12)
<223> Xaa = Cit

<220>
<221> MOD_RES
<222> (14)..(14)
<223> AMIDATION

<400> 17

Xaa Arg Xaa Cys Tyr Xaa Lys Xaa Pro Tyr Arg Xaa Cys Arg
1 5 10

<210> 18
<211> 14
<212> PRT
<213> Artificial Sequence

<220>
<223> Template-fixed peptidomimetic

<220>
<221> MISC_FEATURE
<222> (1)..(1)
<223> Xaa = (PrGU)G

<220>
<221> MISC_FEATURE
<222> (3)..(3)
<223> Xaa = 2-Nal

<220>
<221> DISULFID
<222> (4)..(13)

<220>
<221> MISC_FEATURE
<222> (6)..(6)
<223> Xaa = Cit

<220>
<221> MISC_FEATURE
<222> (8)..(8)
<223> Xaa = D-Pro

<220>
<221> MISC_FEATURE
<222> (12)..(12)

<223> Xaa = Cit

<220>

<221> MOD_RES

<222> (14)..(14)

<223> AMIDATION

<400> 18

Xaa Arg Xaa Cys Tyr Xaa Lys Xaa Pro Tyr Arg Xaa Cys Arg

1 5 10

<210> 19

<211> 14

<212> PRT

<213> Artificial Sequence

<220>

<223> Template-fixed peptidomimetic

<220>

<221> MISC_FEATURE

<222> (1)..(1)

<223> Xaa = (BGU)G

<220>

<221> MISC_FEATURE

<222> (3)..(3)

<223> Xaa = 2-Nal

<220>

<221> DISULFID

<222> (4)..(13)

<220>

<221> MISC_FEATURE

<222> (6)..(6)

<223> Xaa = Cit

<220>

<221> MISC_FEATURE

<222> (8)..(8)

<223> Xaa = D-Pro

<220>

<221> MISC_FEATURE

<222> (12)..(12)

<223> Xaa = Cit

<220>

<221> MOD_RES

<222> (14)..(14)

<223> AMIDATION

<400> 19

Xaa Arg Xaa Cys Tyr Xaa Lys Xaa Pro Tyr Arg Xaa Cys Arg
1 5 10

<210> 20
<211> 14
<212> PRT
<213> Artificial Sequence

<220>
<223> Template-fixed peptidomimetic

<220>
<221> MISC_FEATURE
<222> (3)..(3)
<223> Xaa = 2-Nal

<220>
<221> DISULFID
<222> (4)..(13)

<220>
<221> MISC_FEATURE
<222> (6)..(6)
<223> Xaa = Cit

<220>
<221> MISC_FEATURE
<222> (8)..(8)
<223> Xaa = D-Pro

<220>
<221> MISC_FEATURE
<222> (12)..(12)
<223> Xaa = Cit

<400> 20

Arg Arg Xaa Cys Tyr Xaa Lys Xaa Pro Tyr Arg Xaa Cys Arg
1 5 10

<210> 21
<211> 13
<212> PRT
<213> Artificial Sequence

<220>
<223> Template-fixed peptidomimetic

<220>
<221> MISC_FEATURE
<222> (3)..(3)
<223> Xaa = 2-Nal

<220>
 <221> DISULFID
 <222> (4)..(13)

 <220>
 <221> MISC_FEATURE
 <222> (6)..(6)
 <223> Xaa = Cit

 <220>
 <221> MISC_FEATURE
 <222> (8)..(8)
 <223> Xaa =
 (2S,6S,9S)-6-Amino-2-carboxymethyl-3,8-diazabicyclo-[4,3,0]-nonan
 e-1,4-dione

 <220>
 <221> MISC_FEATURE
 <222> (11)..(11)
 <223> Xaa = Cit

 <220>
 <221> MOD_RES
 <222> (13)..(13)
 <223> AMIDATION

 <400> 21

Arg Arg Xaa Cys Tyr Xaa Lys Xaa Tyr Arg Xaa Cys Arg
 1 5 10

<210> 22
 <211> 13
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Template-fixed peptidomimetic

<220>
 <221> MISC_FEATURE
 <222> (3)..(3)
 <223> Xaa = 2-Nal

<220>
 <221> DISULFID
 <222> (4)..(13)

<220>
 <221> MISC_FEATURE
 <222> (6)..(6)
 <223> Xaa = Cit

<220>

<221> MISC_FEATURE
<222> (8)..(8)
<223> Xaa = AMPA

<220>
<221> MISC_FEATURE
<222> (11)..(11)
<223> Xaa = Cit

<220>
<221> MOD_RES
<222> (13)..(13)
<223> AMIDATION

<400> 22

Arg Arg Xaa Cys Tyr Xaa Lys Xaa Tyr Arg Xaa Cys Arg
1 5 10

<210> 23
<211> 14
<212> PRT
<213> Artificial Sequence

<220>
<223> Template-fixed peptidomimetic

<220>
<221> MISC_FEATURE
<222> (3)..(3)
<223> Xaa = 2-Nal

<220>
<221> DISULFID
<222> (4)..(13)

<220>
<221> MISC_FEATURE
<222> (6)..(6)
<223> Xaa = Cit

<220>
<221> MISC_FEATURE
<222> (9)..(9)
<223> Xaa = D-Pro

<220>
<221> MISC_FEATURE
<222> (12)..(12)
<223> Xaa = Cit

<220>
<221> MOD_RES
<222> (14)..(14)
<223> AMIDATION

<400> 23

Arg Arg Xaa Cys Tyr Xaa Lys Pro Xaa Tyr Arg Xaa Cys Arg
1 5 10

<210> 24

<211> 14

<212> PRT

<213> Artificial Sequence

<220>

<223> Template-fixed peptidomimetic

<220>

<221> MISC_FEATURE

<222> (3)..(3)

<223> Xaa = 2-Nal

<220>

<221> DISULFID

<222> (4)..(13)

<220>

<221> MISC_FEATURE

<222> (6)..(6)

<223> Xaa = Cit

<220>

<221> MISC_FEATURE

<222> (12)..(12)

<223> Xaa = Cit

<220>

<221> MOD_RES

<222> (14)..(14)

<223> AMIDATION

<400> 24

Arg Arg Xaa Cys Tyr Xaa Lys Pro Pro Tyr Arg Xaa Cys Arg
1 5 10

<210> 25

<211> 14

<212> PRT

<213> Artificial Sequence

<220>

<223> Template-fixed peptidomimetic

<220>

<221> MISC_FEATURE

<222> (3)..(3)
<223> Xaa = 2-Nal

<220>
<221> DISULFID
<222> (4)..(13)

<220>
<221> MISC_FEATURE
<222> (6)..(6)
<223> Xaa = Cit

<220>
<221> MISC_FEATURE
<222> (8)..(8)
<223> Xaa = D-Pic

<220>
<221> MISC_FEATURE
<222> (12)..(12)
<223> Xaa = Cit

<220>
<221> MOD_RES
<222> (14)..(14)
<223> AMIDATION

<400> 25

Arg Arg Xaa Cys Tyr Xaa Lys Xaa Pro Tyr Arg Xaa Cys Arg
1 5 10